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\		FIRST NAMED INVENTOR	ATTORNEY DOCKET NO.	CONFIRMATION NO.		
APPLICATION NO.	FILING DATE			5514		
09/535,814	03/28/2000	Yuh-Jiuan Lin	64,600-024 CIP			
	05/06/2002		EXAM	INER		
Tung & Assoc	Tung & Associates					
838 West Long	Lake Road		BRANNOCK, MICHAEL T			
suite 120 Bloomfield Hills, MI 48302			ART UNIT	PAPER NUMBER		
			1646	160		
			DATE MAILED: 05/06/2002	, 14		

Please find below and/or attached an Office communication concerning this application or proceeding.



### UNITED STATES DEPARTMENT OF COMMERCE Patent and Trademark Office COMMISSIONER OF PATENTS AND TRADEMARKS Washington, D.C. 20231

DFA/FCE-1994

SERIAL NUMBER	FILING DATE	FIRST NAMED APPLICANT			
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				E)	XAMINER
•				Michael Brannock	
				ART UNIT	PAPER NUMBER
				1646	16
1			j	DATE MAILED:	

# Please find below a communication from the EXAMINER in charge of this application

Commissioner of Patents

Applicant's response to the Office Action of 3/26/02 is noted, however, the reply filed on 4/22/02 is not fully responsive to the prior Office action because of the following matter(s): this application fails to comply with the requirements of 37 CFR 1.821 through 1.825 as set forth previously. The raw sequence listing is not in a valid format. Please see attached CRF Problem report. Applicant is again encouraged to contact Mark Spencer @ 703 308-4212 for specific help in resolving this continuing problem.

Since the above-mentioned reply appears to be bona fide, applicant is given ONE (1) MONTH or THIRTY (30) DAYS from the mailing date of this notice, whichever is longer, within which to supply the omission or correction in order to avoid abandonment. EXTENSIONS OF THIS TIME PERIOD MAY BE GRANTED UNDER 37 CFR 1.136(a). Extensions of time may be obtained by filing a petition accompanied by the extension fee under the provisions of 37 CFR 1.136(a). In no case may an applicant extend the period for reply beyond the SIX MONTH statutory period. Direct the reply to the undersigned. Applicant is requested to return a copy of the attached Notice to Comply with

Any inquiry concerning this communication or earlier communications from the reply. the examiner should be directed to Michael T. Brannock, Ph.D., whose telephone number is (703) 306-5876. The examiner can normally be reached on Mondays through Fridays from 9:00 a.m. to 5:00 p.m.

If attempts to reach the examiner by telephone are unsuccessful, the

examiner's supervisor, Yvonne Eyler, Ph.D., can be reached at (703) 308-6564

Any inquiry of a general nature or relating to the status of this application or proceeding should be directed to the Group receptionist whose telephone number is (703) 308-0196.

Michael T. Brannock April 30, 2002

SUPERVISORY PATENT EXAMINER TECHNOLOGY CENTER 1600

Application No.: 09535814

NOTICE TO COMPLY WITH REQUIREMENTS FOR PATENT APPLICATIONS CONTAINING NUCLEOTIDE SEQUENCE AND/OR AMINO ACID SEQUENCE DISCLOSURES

The nucleotide and/or amino acid sequence disclosure contained in this application does not comply with the requirements for such a disclosure as set forth in 37 C.F.R. 1.821 - 1.825 for the following reason(s):

X	<ol> <li>This application clearly fails to comply with the requirements of 37 C.F.R. 1.821-1.825. Applicant's attention is directed to these regulations, published at 1114 OG 29, May 15, 1990 and at 55 FR 18230, May 1, 1990.</li> </ol>
	<ol><li>This application does not contain, as a separate part of the disclosure on paper copy, a "Sequence Listing" as required by 37 C.F.R. 1.821(c).</li></ol>
	<ol> <li>A copy of the "Sequence Listing" in computer readable form has not been submitted as required by 37 C.F.R. 1.821(e).</li> </ol>
X	4. A copy of the "Sequence Listing" in computer readable form has been submitted. However, the content of the computer readable form does not comply with the requirements of 37 C.F.R. 1.822 and/or 1.823, as indicated on the attached copy of the marked -up "Raw Sequence Listing."
	5. The computer readable form that has been filed with this application has been found to be damaged and/or unreadable as indicated on the attached CRF Diskette Problem Report. A Substitute computer readable form must be submitted as required by 37 C.F.R. 1.825(d).
	<ol><li>The paper copy of the "Sequence Listing" is not the same as the computer readable from of the "Sequence Listing" as required by 37 C.F.R. 1.821(e).</li></ol>
	7. Other:
Αı	oplicant Must Provide:
X	An <u>initial</u> or substitute computer readable form (CRF) copy of the "Sequence Listing".
X	The trie of the
X	A statement that the content of the paper and computer readable copies are the same and, where applicable, include no new matter, as required by 37 C.F.R. 1.821(e) or 1.821(f) or 1.821(g) or 1.825(b) or 1.825(d).
F	or questions regarding compliance to these requirements, please contact:
F	or Rules Interpretation, call (703) 308-4216
F	or CRF Submission Help, call (703) 308-4212
F	or Patentin software help, call (703) 308-6856

PLEASE RETURN A COPY OF THIS NOTICE WITH YOUR RESPONSE

These pages display submitted file 99/535,8/4C

SEQ ID NOiG1

Met Thr Glu Lys Asn Gln Thr Val Val Ser Glu Phe Val Leu Leu Gly Leu Pro Ile Asp Pro Asp Gln Arg Asp Leu Phe Tyr Ala Leu Phe Leu Ala Met Tyr Val Thr Thr Ile Leu Gly Asn Leu Leu Ile Ile Val Leu Ile Gln Leu Asp Ser His Leu His Thr Pro Met Tyr Leu Phe Leu Ser Asn Leu Ser Phe Ser Asp Leu Cys Phe Ser Ser Val Thr Met Pro Lys Leu Leu Gln Asn Met Gln Ser Gln Val Pro Ser Ile Pro Tyr Ala Gly Cys Leu Thr Gln Met Tyr Phe Phe Leu Phe Phe Gly Asp Leu Glu Ser Phe Leu Leu Val Ala Met Ala Tyr Asp Arg Tyr Val Ala Ile Cys Phe Pro Leu His Tyr Thr Thr Ile Met Ser Pro Lys Leu Cys Phe Ser Leu Leu Val Leu Ser Trp Val Leu Thr Met Phe His Ala Val Leu His Thr Leu Leu Met Ala Arg Leu Cys Phe Cys Ala Asn Thr Ile Pro His Phe Phe Cys Asp Met Ser Ala Leu Leu Lys Leu Ala Cys Ser Asp Thr Gln Val Asn Glu Leu Val Ile Phe Ile Met Gly Gly Leu Ile Leu Val Ile Pro Phe Leu Leu Ile Ile Thr Ser Tyr Ala Arg Ile Val Ser Ser Ile Leu Lys Val Pro Ser Ala Ile Gly Ile Cys Lys Val Phe Ser Thr Cys Gly Ser His Leu Ser Val Val Ser Leu Phe Tyr Gly Thr Val Ile Gly Leu Tyr Leu Cys Pro Ser Ala Asn Asn Ser Thr Val Lys Glu Thr Ile Met Ala Met Met Tyr Thr Val Val Thr Pro Met Leu Asn Pro Phe Ile Tyr Ser Leu Arg Asn Lys Asp Met Lys Gly Ala Leu Arg Arg Val Ile Cys Arg Lys Lys Ile Thr Phe Ser Val

SEQ ID NOiG2 Asp-Pro-Asp-Gln-Arg-Asp-Cys

SEQ ID NOiG3 Leu-Phe-Leu-Ser-Asn-Leu-Ser-Phe-Ser-Asp-Leu-Cys-Ala

The file submitted is hot a Sequence Listing. Cleare see sample Sequence Listing (attacked in back) for valid format. Also, please consult Sequene Rules,

SEQUENCE LISTING

SEQ ID NOIG1 LENGTHíG313 ORGANISMíGCanis familiaris SEQUENCEiGP30955

DBSOURCEiGswissprot: locus OLFD\_CANFA, accession P30955

@/535,814C ]

SEQ ID NOiG2 LENGTHiG7 TYPEiGPRT ORGANISMiGCanis familiaris SEQUENCEiGB1 Asp-Pro-Asp-Gln-Arg-Asp-Cys

SEQ ID NOiG3 LENGTHiG13 TYPEiGPRT ORGANISMiGCanis familiaris SEQUENCEiGPb2 Leu-Phe-Leu-Ser-Asn-Leu-Ser-Phe-Ser-Asp-Leu-Cys-Ala

# Please consult.

<110>	Smith, John: Smithgene Inc.
<120>	Example of a Sequence Listing
<130>	01-00001
<140> <141>	PCT/EP98/00001 1998-12-31
<150> <151>	US 08/999.999 1997-10-15
<160>	
<170>	PatentIn version 2.0
<210 > <211 > <211 > <212 > <213 > <220 > <221 > <221 > <221 > <222 > <300 > <301 > <302 > <	1 389 DNA Paramecium sp.  CDS (279)(389)  Doe. Richard Isolation and Characterization of a Gene Encoding a Protease (rom Paramecium sp. Journal of Genes
<303> <304> <305> <306> <307> <308> <308> <309>	1 4 1 - 7 1 988 - 06 - 31 1 2 3 4 5 6 1 9 8 8 - 06 - 31
<400> agciglagic	l acceptot ceretrete ergggerici caccetgera areagatete 60
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rgargragca	attgctggca gtgccacagg cttttcagcc aggcttaggg tgggttccgc 180
cacaacacaa	eggeeetet egegeteete tegegeetet etetegetet eetetegete 240

### Appendix 3, page 2

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ttc
                                                                                        AQC
                                                                                               296
                                                                 gtt
                                                                       tca
                                                                             atq
                                                           atg
                                           cagttage
                            9499499999
                                                                                   Phe
                                                                                        Ser
              aggtgagcag
                                                                 Val
                                                                       Ser
                                                                             Mct
ggacctgatt
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                                                          ctg cag ccg aat ctt
Leu Gln Pro Asn biu
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      Pro
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                                    Lcu
                 Het Phe
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      Val
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      val
            Cys
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                   20
                       Lou
            Pro Asn
      Cln
Leu
             35⊕
<210>
<211>
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              rrt
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              Artificial Sequence
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              Designed pentide based on size and polarity to act as a
linker between the alpha and beta chains of Protein XYZ.
<220>
<223>
<400>
                              tro Met His Thr Glu
                                                          He
                       Clu
Met Val
            Λsn
                 Leu
                                                      10
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<210>
<400>
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[Annex VIII follows]

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table. The numeric identifier shall be used only in the Sequence Listing." The order and presentation of the items of information in the "Sequence Listing" shall conform to the arrangement given below. Each "Sequence Listing" shall conform to the arrangement given below. Each item of information shall begin on a new line and shall begin with the item of information shall begin on a new line and shall begin with the numeric identifier enclosed in angle brackets as shown. The submission of those items of information designated with an "M" is mandatory. The submission of those items of information designated with an "O" is submission of those items of information designated with an "O" is optional. Numeric identifiers <110> through <170> shall only be set forth at the beginning of the "Sequence Listing." The following table illustrates the numeric identifiers.

illustrates	the numeric identi	fiers.	•
Numeric Identifier	Definition	Comments and Format	Mandatory (M) or Optional (O)
<110>	Applicant	Preferably max. of 10 names; one name per line; preferable format: Surname, Other: Names and/or Initials	M v
<120>	Title of Invention		<b>н</b>
<130>	File Reference	Personal file reference	M, when filed prior to assignment of appl. number
<140>	Current Applica- tion Number	Specify as: US 07/999,999 or PCT/US96/99999	M, if available
<141>	Current Filing Date	Specify as: yyyy-mm-dd	•
<150>	Prior Application Number	Specify as: US 07/999,999 or PCT/US96/99999	M, if applicable include priority documents under 35 USC 119 and 120
<151>	Prior Application Filing Date	Specify as: yyyy-mm-dd	M, i[ applicable
<160>	Number of SEQ ID	Count includes total number of SEQ ID NOs	М
<170>	Software	Name of software used to create the Sequence Listing	0
<210>	SEQ ID NO: #:	Response shall be an integer repressenting the SEQ ID NO shown	M .
<211>	Length	Respond with an integer expressing the number of bases or amino acid residues	м .

... Whether presented sequence molecule is DNA, RNA, or PRT (protein). If a nucleotide sequence contains both DNA and RNA fragments, the type shall be "DNA." In addition, the combined DNA/ RNA molecule shall be further described in . the <220> to <223> [cature section.

<213>

Organism -

Scientific name, i.e. Genus/species, Unknown or Artificial Sequence. In addition, the "Unknown" or "Artificial Sequence" organisms shall be further described in the <220> to <223> feature section.

<220>

Feature

Leave blank after <220>. <221-223> provide for a description of points of biological significance in the sequence.

M, under the following conditions: if "n,"
"Xaa," or a modified or unusual L-amino acid or modified base was used in a sequence; if ORGAN-ISM is "Artificial Sequence" or "Unknown"; if molecule is combined DNA/RNA.

<221>

Name/Key

Provide appropriate identifier for feature, preferably from WIPO Standard ST.25 (1998), Appendix 2, Tables 5 and 6

M, under the following conditions: if "n," "Xaa," or a modified or unusual L-amino acid or modified base was used in a sequence

<222>

Location

Specify location within sequence; where appropriate state number of first and last bases/amino acids

M, under the following conditions: if "n," "Xaa," or a modified or unusual L-amino acid or modified

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e was used in sequence

**:** . . M, under the fol-Other relevant Other Inforlowing conditi ns: <223> information; if "n," "Xaa," or mation "four lines maximum a modified or unusual L-amino acid or modified base was used in a sequence; if **ORGANISM** is "Artificial Sequence" or "Unknown"; .i.(\_= molecule is combined DNA/RNA-Leave blank Publication <300> after <300> Information Preferably max **Authors** <301> of ten named authors of publication; specify one name per line; preferable format: Surname, Other Names and/or Initials ġ Title <302> 0 Journal <303> 0 Volume <304> 0 Issuc <305> ο. Pages <306> Journal date on which O <307> Date data published; specify as yyyy-mmdd, 100M-yyyy or Scason-yyyy 0 Accession number <300> Database assigned by data-Accession: base including Number . database name

Date of entry in

as yyyy-mm-dd or

Document number;

for patent-type

citations only. Specify as, for example, US 07/999.999

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Database Entry

Patent Document

Date:

Number

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<312>	Publication Date	Document publication date, for patent-type citations only; specify as yyyy-mm-do		•
		•	•	•
<313>	Relevant Residues	FROM (position) TO (position)	0	ب الأحسا
<400>		SEQ ID NO should follow the numeric identifier and should appear on the line preceding the actual sequence	M V	
	*	acqueilee /	· 🗸	

- 5. Section 1.824 is revised to read as follows:
- 1.824 Form and format for nucleotide and/or amino acid sequence submissions in computer readable form.
- (a) The computer readable (orm required by 1.821(c) shall meet the following specifications:
- (1) The computer readable (orm shall contain a single "Sequence Listing" as either a diskette, series of diskettes, or other permissible media outlined in paragraph (c) of this section.
- (2) The "Sequence Listing" in paragraph (a) (1) of this section shall be submitted in American Standard Code for Information Interchange (ASCII) text. No other formats shall be allowed.
- (3) The computer readable form may be created by any means, such as word processors, nucleotide/amino acid sequence editors or other custom computer programs; however, it shall conform to all specifications detailed in this section.
- (4) File compression is acceptable when using diskette media, so long as the compressed file is in a self-extracting format that will decompress on one of the systems described in paragraph (b) of this section.
- (5) Page numbering shall not appear within the computer readable form version of the "Sequence Listing" (ile.
- (6) All computer readable (orms shall have a label permanently affixed thereto on which has been hand-printed or typed: the name of the applicant, the title of the invention, the date on which the data were recorded on the computer readable form, the operating system used, a reference number, and an application serial number and filing date, if known.
- (b) Computer readable (orm submissions must meet these format requirements:
- (1) Computer: IBM PC/XT/AT, or compatibles, or Apple Macintosh;
- (2) Operating System: MS-DOS, Unix or Macintosh:

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